



1600

RAW SEQUENCE LISTING

DATE: 11/26/2002

PATENT APPLICATION: US/09/581,976B

TIME: 09:26:10

Input Set : A:\seqlist.txt

Output Set: N:\CRF4\11262002\I581976B.raw

ENTERED

4 <110> APPLICANT: Dalemans, Wilfried L.J.
 5 Gerard, Catherine Marie Ghislaine
 7 <120> TITLE OF INVENTION: Vaccine
 10 <130> FILE REFERENCE: B45124
 12 <140> CURRENT APPLICATION NUMBER: 09/581,976B
 13 <141> CURRENT FILING DATE: 2000-06-20
 15 <150> PRIOR APPLICATION NUMBER: PCT/EP98/08563
 16 <151> PRIOR FILING DATE: 1998-12-18
 18 <150> PRIOR APPLICATION NUMBER: GB 9727262.9
 19 <151> PRIOR FILING DATE: 1997-12-24
 21 <160> NUMBER OF SEQ ID NOS: 28
 23 <170> SOFTWARE: FastSEQ for Windows Version 3.0
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 26 <211> LENGTH: 220
 27 <212> TYPE: PRT
 28 <213> ORGANISM: Artificial Sequence
 30 <220> FEATURE:
 31 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
 32 influenza B and E7 from Human papilloma virus type
 33 16)
 35 <400> SEQUENCE: 1
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 37 1 5 10 15
 38 Ser Asp Lys Ile Ile Ile Ala His Arg Gly Ala Ser Gly Tyr Leu Pro
 39 20 25 30
 40 Glu His Thr Leu Glu Ser Lys Ala Leu Ala Phe Ala Gln Gln Ala Asp
 41 35 40 45
 42 Tyr Leu Glu Gln Asp Leu Ala Met Thr Lys Asp Gly Arg Leu Val Val
 43 50 55 60
 44 Ile His Asp His Phe Leu Asp Gly Leu Thr Asp Val Ala Lys Lys Phe
 45 65 70 75 80
 46 Pro His Arg His Arg Lys Asp Gly Arg Tyr Tyr Val Ile Asp Phe Thr
 47 85 90 95
 48 Leu Lys Glu Ile Gln Ser Leu Glu Met Thr Glu Asn Phe Glu Thr Met
 49 100 105 110
 50 Ala Met His Gly Asp Thr Pro Thr Leu His Glu Tyr Met Leu Asp Leu
 51 115 120 125
 52 Gln Pro Glu Thr Thr Asp Leu Tyr Cys Tyr Glu Gln Leu Asn Asp Ser
 53 130 135 140
 54 Ser Glu Glu Glu Asp Glu Ile Asp Gly Pro Ala Gly Gln Ala Glu Pro
 55 145 150 155 160
 56 Asp Arg Ala His Tyr Asn Ile Val Thr Phe Cys Cys Lys Cys Asp Ser
 57 165 170 175

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58 Thr Leu Arg Leu Cys Val Gln Ser Thr His Val Asp Ile Arg Thr Leu
59          180          185          190
60 Glu Asp Leu Leu Met Gly Thr Leu Gly Ile Val Cys Pro Ile Cys Ser
61          195          200          205
62 Gln Lys Pro Thr Ser Gly His His His His His
63          210          215          220
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66 <211> LENGTH: 663
67 <212> TYPE: DNA
68 <213> ORGANISM: Artificial Sequence
70 <220> FEATURE:
71 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
72     influenza B and E7 from Human papilloma virus type
73     16)
75 <400> SEQUENCE: 2
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77 attattgctc accgtggtgc tagcggttat ttaccagagc atacgttaga atctaaagca      120
78 cttgcgtttg cacaacaggc tgattattta gagcaagatt tagcaatgac taaggatggt      180
79 cgttttagtg ttattcacga tcacttttta gatggcttga ctgatgttgc gaaaaaattc      240
80 ccacatcgtc atcgtaaaga tggccgttac tatgtcatcg actttacctt aaaagaaatt      300
81 caaagttag aaatgacaga aaactttgaa accatggcca tgcattgaga tacacctaca      360
82 ttgcatgaat atatgttaga ttgtcaacca gagacaactg atctctactg ttatgagcaa      420
83 ttaaagtaca gtcagagga ggaggatgaa atagatggtc cagctggaca agcagaaccg      480
84 gacagagccc attacaatat tgtaaccttt tgttgcaagt gtgactctac gcttcggttg      540
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92 <213> ORGANISM: Artificial Sequence
94 <220> FEATURE:
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97     16)
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101 attattgctc accgtggtgc tagcggttat ttaccagagc atacgttaga atctaaagca      120
102 cttgcgtttg cacaacaggc tgattattta gagcaagatt tagcaatgac taaggatggt      180
103 cgttttagtg ttattcacga tcacttttta gatggcttga ctgatgttgc gaaaaaattc      240
104 ccacatcgtc atcgtaaaga tggccgttac tatgtcatcg actttacctt aaaagaaatt      300
105 caaagttag aaatgacaga aaactttgaa accatggcca tgtttcagga cccacaggag      360
106 cgaccagaa agttaccaca gttatgcaca gagctgcaaa caactataca tgatataata      420
107 ttagaatgtg tgtactgcaa gcaacagtta ctgacgacgtg aggtatatga ctttgctttt      480
108 cgggatttat gcatagtata tagagatggg aatccatatg ctgtatgtga taaatgttta      540
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110 ttagaacagc aatacaaaa accgttgtgt gatttggtta ttaggtgtat taactgtcaa      660
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112 ataaggggtc ggtggaccgg tcgatgtatg tcttggttga gatcatcaag aacacgtaga      780

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121 <220> FEATURE:
122 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
123     influenza B and E6 from Human papilloma virus type
124     16)
126 <400> SEQUENCE: 4
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129 Ser Asp Lys Ile Ile Ile Ala His Arg Gly Ala Ser Gly Tyr Leu Pro
130          20          25          30
131 Glu His Thr Leu Glu Ser Lys Ala Leu Ala Phe Ala Gln Gln Ala Asp
132          35          40          45
133 Tyr Leu Glu Gln Asp Leu Ala Met Thr Lys Asp Gly Arg Leu Val Val
134          50          55          60
135 Ile His Asp His Phe Leu Asp Gly Leu Thr Asp Val Ala Lys Lys Phe
136          65          70          75          80
137 Pro His Arg His Arg Lys Asp Gly Arg Tyr Tyr Val Ile Asp Phe Thr
138          85          90          95
139 Leu Lys Glu Ile Gln Ser Leu Glu Met Thr Glu Asn Phe Glu Thr Met
140          100         105         110
141 Ala Met Phe Gln Asp Pro Gln Glu Arg Pro Arg Lys Leu Pro Gln Leu
142          115         120         125
143 Cys Thr Glu Leu Gln Thr Thr Ile His Asp Ile Ile Leu Glu Cys Val
144          130         135         140
145 Tyr Cys Lys Gln Gln Leu Leu Arg Arg Glu Val Tyr Asp Phe Ala Phe
146          145         150         155         160
147 Arg Asp Leu Cys Ile Val Tyr Arg Asp Gly Asn Pro Tyr Ala Val Cys
148          165         170         175
149 Asp Lys Cys Leu Lys Phe Tyr Ser Lys Ile Ser Glu Tyr Arg His Tyr
150          180         185         190
151 Cys Tyr Ser Leu Tyr Gly Thr Thr Leu Glu Gln Gln Tyr Asn Lys Pro
152          195         200         205
153 Leu Cys Asp Leu Leu Ile Arg Cys Ile Asn Cys Gln Lys Pro Leu Cys
154          210         215         220
155 Pro Glu Glu Lys Gln Arg His Leu Asp Lys Lys Gln Arg Phe His Asn
156          225         230         235         240
157 Ile Arg Gly Arg Trp Thr Gly Arg Cys Met Ser Cys Cys Arg Ser Ser
158          245         250         255
159 Arg Thr Arg Arg Glu Thr Gln Leu Thr Ser Gly His His His His His
160          260         265         270
161 His
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165 <211> LENGTH: 1116
166 <212> TYPE: DNA
167 <213> ORGANISM: Artificial Sequence

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169 <220> FEATURE:
170 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
171     influenza B and E6E7 fusion from Human papilloma
172     virus type 16)
174 <400> SEQUENCE: 5
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177 cttgcgtttg cacaacaggc tgattattta gagcaagatt tagcaatgac taaggatggt      180
178 cgttttagtg ttattcacga tcacttttta gatggcttga ctgatgttgc gaaaaaattc      240
179 ccacatcgtc atcgtaaaga tggccgttac tatgtcatcg actttacctt aaaagaaatt      300
180 caaagtttag aaatgacaga aaactttgaa accatggcca tgtttcagga cccacaggag      360
181 cgaccagaaa agttaccaca gttatgcaca gagctgcaaa caactataca tgatataata      420
182 ttagaatgtg tgtactgcaa gcaacagtta ctgcgacgtg aggtatatga ctttgctttt      480
183 cgggatttat gcatagtata tagagatggg aatccatatg ctgtatgtga taaatgttta      540
184 aagttttatt ctaaaattag tgagtataga cattattggt atagtttgta tggaacaaca      600
185 ttagaacagc aatacaacaa accgttgtgt gatttgtaa ttaggtgtat taactgtcaa      660
186 aagccactgt gtcctgaaga aaagcaaaga catctggaca aaaagcaaag attccataat      720
187 ataaggggtc ggtggaccgg tcgatgtatg tcttggtgca gatcatcaag aacacgtaga      780
188 gaaacccagc tgatgcatgg agatacacct acattgcatg aatatatggt agatttgcaa      840
189 ccagagacaa ctgatctcta ctgttatgag caattaaatg acagctcaga ggaggaggat      900
190 gaaatagatg gtccagctgg acaagcagaa ccggacagag cccattacaa tattgtaacc      960
191 ttttggtgca agtgtgactc tacgcttcgg ttgtgcgtac aaagcacaca cgtagacatt     1020
192 cgtactttgg aagacctgtt aatgggcaca ctagggaattg tgtgccccat ctgttctcag     1080
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196 <211> LENGTH: 371
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202     influenza B and E6E7 fusion from Human papilloma
203     virus type 16)
205 <400> SEQUENCE: 6
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209           20               25               30
210 Glu His Thr Leu Glu Ser Lys Ala Leu Ala Phe Ala Gln Gln Ala Asp
211           35               40               45
212 Tyr Leu Glu Gln Asp Leu Ala Met Thr Lys Asp Gly Arg Leu Val Val
213           50               55               60
214 Ile His Asp His Phe Leu Asp Gly Leu Thr Asp Val Ala Lys Lys Phe
215           65               70               75               80
216 Pro His Arg His Arg Lys Asp Gly Arg Tyr Tyr Val Ile Asp Phe Thr
217           85               90               95
218 Leu Lys Glu Ile Gln Ser Leu Glu Met Thr Glu Asn Phe Glu Thr Met
219           100              105              110
220 Ala Met Phe Gln Asp Pro Gln Glu Arg Pro Arg Lys Leu Pro Gln Leu
221           115              120              125

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222 Cys Thr Glu Leu Gln Thr Thr Ile His Asp Ile Ile Leu Glu Cys Val
223      130                      135                      140
224 Tyr Cys Lys Gln Gln Leu Leu Arg Arg Glu Val Tyr Asp Phe Ala Phe
225      145                      150                      155                      160
226 Arg Asp Leu Cys Ile Val Tyr Arg Asp Gly Asn Pro Tyr Ala Val Cys
227      165                      170                      175
228 Asp Lys Cys Leu Lys Phe Tyr Ser Lys Ile Ser Glu Tyr Arg His Tyr
229      180                      185                      190
230 Cys Tyr Ser Leu Tyr Gly Thr Thr Leu Glu Gln Gln Tyr Asn Lys Pro
231      195                      200                      205
232 Leu Cys Asp Leu Leu Ile Arg Cys Ile Asn Cys Gln Lys Pro Leu Cys
233      210                      215                      220
234 Pro Glu Glu Lys Gln Arg His Leu Asp Lys Lys Gln Arg Phe His Asn
235      225                      230                      235                      240
236 Ile Arg Gly Arg Trp Thr Gly Arg Cys Met Ser Cys Cys Arg Ser Ser
237      245                      250                      255
238 Arg Thr Arg Arg Glu Thr Gln Leu Met His Gly Asp Thr Pro Thr Leu
239      260                      265                      270
240 His Glu Tyr Met Leu Asp Leu Gln Pro Glu Thr Thr Asp Leu Tyr Cys
241      275                      280                      285
242 Tyr Glu Gln Leu Asn Asp Ser Ser Glu Glu Glu Asp Glu Ile Asp Gly
243      290                      295                      300
244 Pro Ala Gly Gln Ala Glu Pro Asp Arg Ala His Tyr Asn Ile Val Thr
245      305                      310                      315                      320
246 Phe Cys Cys Lys Cys Asp Ser Thr Leu Arg Leu Cys Val Gln Ser Thr
247      325                      330                      335
248 His Val Asp Ile Arg Thr Leu Glu Asp Leu Leu Met Gly Thr Leu Gly
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250 Ile Val Cys Pro Ile Cys Ser Gln Lys Pro Thr Ser Gly His His His
251      355                      360                      365
252 His His His
253      370
255 <210> SEQ ID NO: 7
256 <211> LENGTH: 663
257 <212> TYPE: DNA
258 <213> ORGANISM: Artificial Sequence
260 <220> FEATURE:
261 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
262      influenza B and mutated E7 from Human papilloma
263      virus type 16)
265 <400> SEQUENCE: 7
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267 attattgctc accgtggtgc tagcggttat ttaccagagc atacgtaga atctaaagca      120
268 cttgcgtttg cacaacaggc tgattattta gagcaagatt tagcaatgac taaggatggt      180
269 cgttttagtgg ttattcacga tcacttttta gatggcttga ctgatgttgc gaaaaaattc      240
270 ccacatcgtc atcgtaaaga tggccgttac tatgtcatcg actttacctt aaaagaaatt      300
271 caaagtttag aaatgacaga aaactttgaa accatggcca tgcattgaga tacacctaca      360
272 ttgcatgaat atatgttaga ttgcaacca gagacaactg atctctacgg ttatcagcaa      420
273 ttaaattgaca gctcagagga ggaggatgaa atagatggtc cagctggaca agcagaaccg      480

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VERIFICATION SUMMARY

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